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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant(s): LI et al.

Appln No.: 10/075,105

Filed: February 13, 2002

For: A BI-DIRECTIONAL DUAL PROMOTER
COMPLEX WITH ENHANCED
PROMOTER ACTIVITY FOR TRANSGENE
EXPRESSION IN EUKARYOTES

Group Art

Unit: 1645

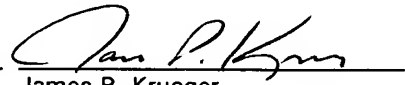
Examiner: Not Yet Known

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Date



James P. Krueger

Registration No. 35, 234

Attorney for Applicant(s)

TRANSMITTAL OF FORMAL DRAWINGS

Box MISSING PARTS

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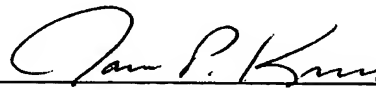
ATTENTION: Assistant Commissioner for Patents

Washington, D.C. 20231

Sir:

To correct informalities in the drawings as noted in the Notice to File Missing Parts of Nonprovisional Application dated April 18, 2002, Applicants submit herewith formal drawings (Figure 1 -27 on 37 sheets) for this application. The two-month period of response set in the Notice of Missing Parts expired on June 18, 2002, as a result this submission includes a two-month Petition for Extension of Time.

Respectfully submitted,
FITCH, EVEN, TABIN & FLANNERY

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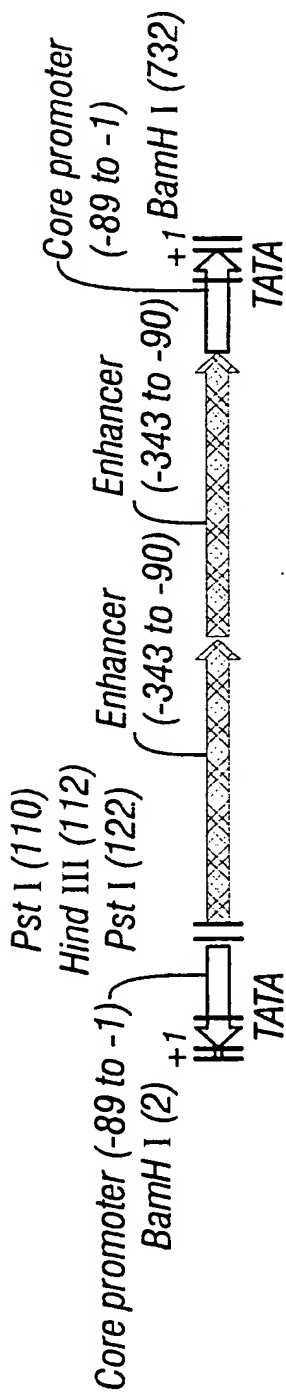


FIG. 1

BamHI

1 GGATCCAGCG TGTCTCTCICC AAATGAAATG AACTTCCTTA TATAGAGGAA GGGTCTTGCG AAGGATAGTG GGATTGTGCG
CCTAGGTGCG ACAGGAGAGG TTTACTTTAC TTGAAGGAAT ATAATCTCTT CCCAGAAGCG TTCCTATCAC CCTAACACGC

PstI HindIII PstI

81 TCATCCCTTA CGTCAGTGGG GATACTGCAG AAGCTTCTGC AGTGAGACTT TTCAACAAAG GGTAAATATCG GGAACCTICC
AGTAGGGAAT GCAGTCACCT CTATGACGTC TTCGAAGACG TCACTCTGAA AAGTTGTTTC CCATTATAGC CCTTTGGAGG

161 TCGGATTCCA TTGCCCAGCT ATCTGTCACT TCATCAAAAG GACAGTAGAA AAGGAAGGTG GCACCTACAA ATGCCATCAT
AGCCTAAGGT AACGGGTGCGA TAGACAGTGA AGTAGTTTC CTGTCTCTT TTCCTTCCAC CGTGGATGTT TACGGTAGTA

241 TCGGATAAAG GAAAGGCTAT CGTTCAAGAT GCCTCTGCCG ACAGTGGTCC CAAAGATGGA CCCCACCCCA CGAGGAGCAT
ACGCTATTTC CTTTCGGATA GCAAGTTCTA CGGAGACGGC TGTCACCAGG GTTCTACCT GGGGGTGGT GCTCCTCGTA

321 CGTGGAAGAA GAAGACGTTT CAACCACGTC TTCAAGCAA GTGGATTGAT GTGATTGCAG TGAGACTTTT CAACAAAGGG
GCACCTTTT CTTCTGCAAG GTTGGTGCAG AAGTTTCGT CACCTAATA CACTAACGTC ACTCTGAAA GTTGTTTCCC

401 TAATATCGGG AAACCTCCTC GGATTCCATT GCCCAGCTAT CTGTCACTTC ATCAAAAGGA CAGTAGAAAA GGAAGGTGGC
ATTATAGCCC TTGGAGGAG CTAAGGTAA CGGGTCGATA GACAGTGAAG TAGTTTCTCT GTCACTTTT CCTTCCACCG

481 ACCTACAAAT GCCATCATTG CGATAAAGGA AAGGCTATCG TTCAAGATGC CTCTGCCGAC AGTGTGCCCA AAGATGGACC
TGGATGTTTA CGGTAGTAAC GCTATTTCCT TTCCGATAGC AAGTTCTACG GAGACGGCTG TCACCAGGTT TTCTACCTGG

FIG. 2A

561 CCCACCCACG AGGAGCATCG TGGAAAAGA AGACGTTCCA ACCACGTCTT CAAAGCAAGT GGATTGATGT GATATCTCCA
 GGGTGGGTGC TCCTCGTAGC ACCTTTTCT TCTGCAAGT TGGTGCAGAA GTTTCGTTCA CCTAACTACA CTATAGAGGT

641 CTGACGTAAG GGATGACGCA CAATCCCACT ATCCTTCGCA AGACCCCTCC TCTATATAAG GAAGTTCATT TCATTGGAG
 GACTGCATTG CCTACTGCGT GTTAGGGTGA TAGGAAGCGT TCTGGGAAGG AGATATATTC CTTCAAGTAA AGTAAACCTC

BamHI

721 AGGACACGCT GGATCC Seq. ID No. 1
 TCCTGTGCGA CCTAGG Seq. ID No. 2

FIG. 2B

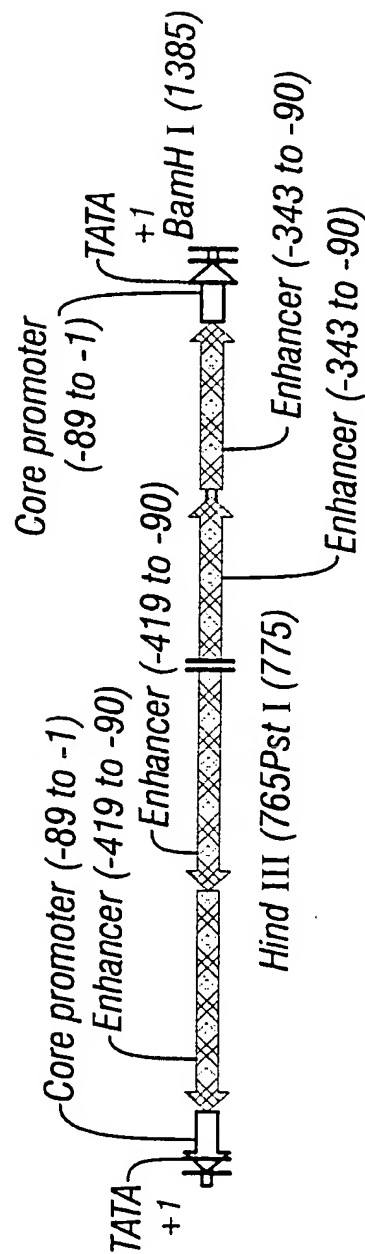


FIG. 3



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SnaBI

Seq. ID No. 3

1 TAGGTACAGC GTGTCCTCTC CAAATGAAT GAACCTCCTT ATATAGAGGA AGGTCCTTGC GAAGGATAGT GGGATTGTGC

Seq. ID No. 4

ATGCATGTGC CACAGGAGAG GTTTACTTTA CTTGAAGGAA TATATCTCCT TCCAGAAGC CTTCTATCA CCCTAACACG

81 GTCATCCCTT ACGTCAGTGG AGATATCACA TCCATCCACT TGCCTTGAAG ACGTGGTTGG AACGTCTTCT TTTCCACGA
CAGTAGGGAA TGCAGTCACC TCTATAGTGT AGTTAGGTGA ACGAACTTC TGCACCAACC TTGCAGAAGA AAAAGGTGCT

161 TGCTCCTCGT GGGTGGGGT CCATCTTTGG GACCACTGTC GGCAGAGGCA TCTTCAACGA TGGCCTTTCC TTTATCGCAA
ACGAGGAGCA CCCACCCCA GGTAGAAACC CTGGTGACAG CCGTCTCCGT AGAAGTTGCT ACCGGAAGG AAATAGCGTT

241 TGATGGCAAT TGTAGGAGCC ACCTTCCTTT TCCACTATCT TCACAATAAA GTGACAGATA GCTGGGCAAT GGAATCCGAG
ACTACCGTAA ACATCCTCGG TGAAGGAAA AGGIGATAGA AGTGTTATTT CACTGTCTAT CGACCCGTTA CCTTAGGCTC

321 GAGGTTTCCG GATATTACCC TTTGTTGAAA AGTCTCAATT GCCCTTTGGT CTTCTGAGAC TGTATCTTTG ATATTTTGG
CTCCAAGGC CTATAATGGG AAACAACCTT TCAGAGTTAA CGGGAACCA GAAGACTCTG ACATAGAAAC TATAAAACC

401 AGTAGACAAG TGTGTCGTGC TCCACCATGT TGATTCACAT CAATCCACTT GCTTTGAAGA CGTGGTTGA ACGTCTTCTT
TCATCTGTTC ACACAGCAGC AGGTGGTACA ACTAAGTGA GTTAGGTGAA CGAACTTCT GCACCAACCT TGCAGAAGAA

481 TTTCCACGAT GCTCCTCGTG GGTGGGGGTC CATCTTTGGG ACCACTGTG GCAGAGGCAT CTTCAACGAT GGCCTTTCCT
AAAGGTGCTA CGAGGAGCAC CCACCCCCAG GTAGAAACCC TGGTGACAGC CGTCTCCGTA GAAGTTGCTA CCGGAAGGA

FIG. 4A

561 TTATCGCAAT GATGGCATT GTAGGAGCCA CCTTCCTTT CCACIACTT CACAATAAG TGACAGATAG CTGGGCAATG
AATAGCGTTA CTACCGTAAA CATCTCGGT GGAAGGAAA GTGATAGAA GTGTTAATTC ACTGTCTATC GACCCGTTAC

641 GAATCCGAGG AGGTTTCGG ATATTACCCT TTGTTGAAA GTICAAATG CCCTTGGTC TTCTGAGACT GTATCTTGA
CTTAGGCTCC TCCAAAGGCC TATAATGGGA AACAACCTTT CAGAGTTAAC GGGAAACCAG AAGACTCTGA CATAGAAACT

HindIIIIPstI

721 TATTTTIGGA GTAGACAAAGT GTGTCGTGCT CCACCATGTT GATAAGCTTC TGCAGTGAGA CTTTCAACA AAGGTAATA
ATAAAACCT CATCTGTTCA CACAGCACGA GTGGGTACAA CTATTCGAAG ACGTCACICT GAAAAGTTGT TTCCCATTA

801 TCGGGAACC TCCTCGGATT CCATTGCCCA GCTATCTGTC ACTTCATCAA AAGGACAGTA GAAAAGGAAG GTGGCACCTA
AGCCCTTTGG AGGAGCCTAA GGTAACGGGT CGATAGACAG TGAAGTAGT TTCTGTICAT CTTTTCCTTC CACCGTGGAT

881 CAAATGCCAT CATTGCGATA AAGGAAAGGC TATCGTTCAA GATGCCTCTG CCGACAGTGG TCCCAAGAT GGACCCCCAC
GTTTACGGTA GTAACGCTAT TTCTTTCCG ATAGCAAGT CTACGGAGAC GGCTGTCACC AGGTTTCTA CCTGGGGGTG

961 CCACGAGGAG CATCGTGGAA AAAGAAGACG TTCCAACCAC GTCTTCAAAG CAAGTGGATT GATGTATTG CAGTGAGACT
GGTGCTCCTC GTAGCACCTT TTCTTCTGC AAGGTGGTG CAGAAGTTT GTTCACCTAA CTACACTAAC GTCACCTIGA

1041 TTCAACAAA GGGTAATAIC GGGAAACCTC CTCGGATTCC ATTGCCCAGC TATCTGTCAC TTCATCAAAA GGACAGTAGA
AAAGTTGTT CCCATTATAG CCCTTTGGAG GAGCCTAAGG TAACGGGTG ATAGACAGTG AAGTAGTTTT CCTGTCAICT

1121 AAAGGAAGGT GGCACCTACA AATGCCATCA TTGGGATAA GGAAAGGCTA TCGTTCAAGA TGCCTCTGCC GACAGTGGTC
TTTCTTCCA CCGTGGATGT TTACGGTAGT AAGCTATTT CCTTCCGAT AGCAAGTTCT ACGGAGACGG CTGTCACCAG

FIG. 4B

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1201 CCAAAGATGG ACCCCACCCC ACGAGGAGCA TCGTGGA AAA AGAAGACGTT CCAACCACGT CTCCTAAGCA AGTGGATTGA
 GGTTCCTACC TGGGGGTGGG TGCCTCCTCGT AGCACCCTTT TCTTCTGCAA GGTGGTGCA GAAGTTTCGT TCACCTAACT
 1281 TGTGATATCT CCACTGACGT AAGGGATGAC GCACAATCCC ACTATCCTTC GCAAGACCCT TCCTCTATAT AAGGAAGTTC
 AACTATAGA GGTGACTGCA TTCCTACTG CGTGTAGGG TGATAGGAAG CGTCTGGGA AGGAGATATA TTCCTTCAAG

FIG. 4C

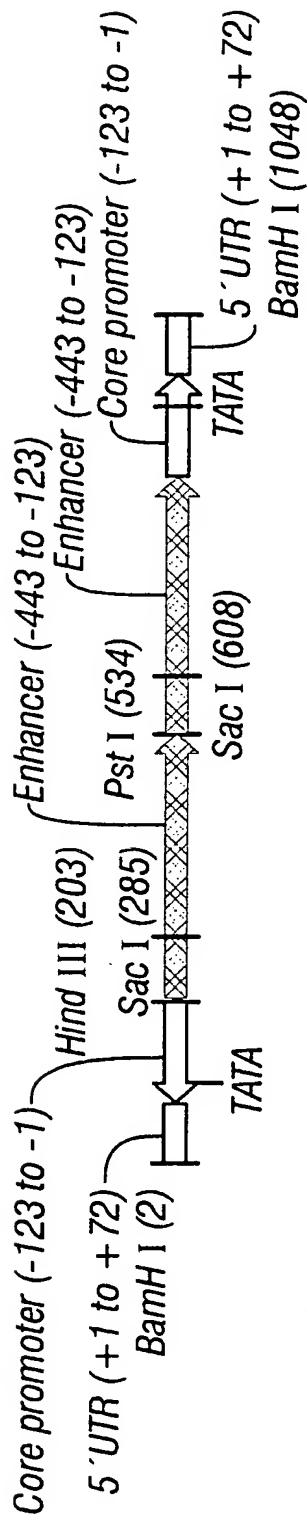


FIG. 5

BamHI

1 GGATCCACAA ACTTACAAAT TTCTCTGAAG TTGTATCCTC AGTACTTCAA AGAAAATAGC TTACACCAA TTTTTCCTTG
CCTAGGTGTT TGAATGTTTA AAGAGACTTC AACATAGGAG TCATGAAGTT TCTTTTATCG AATGTGGTTT AAAAAAGAAC
81 TTTTCACAAA TGCCGAACCT GGTTCCTTAT ATAGGAAAAC TCAAGGGCAA AAATGACACG GAAAAATATA AAAGGATAAG
AAAAGTGTTT ACGGCTTGAA CCAAGGAATA TATCCTTTTG AGTTCCCGTT TTTACTGTGC CTTTTTATAT TTTCCTATTG

HindIII

161 TAGTGGGGGA TAAGATTCTT TTGTGATAAG GTTACTTTCC GAAGCTTCCA GAAGGTAATT ATCCAAGATG TAGCATCAAG
ATCACCCCCT ATTCTAAGGA AACACTATTG CAATGAAAGG CTTGGAAGGT CTTCCATTAA TAGGTTCTAC ATCGTAGTTC

SacI

241 AATCCCAATGT TTACGGGAAA AACTATGGAA GTATTATGTG AGCTCAGCAA GAAGCAGATC AATATGCGGC ACATATGCAA
TTAGGTACA AATGCCCTTT TTGATACCTT CATAATACAC TCGAGTCGTT CTTGCTCTAG TTATACGCCG TGTATACGTT
321 CCTATGTTCA AAAATGAAGA ATGTACAGAT ACAAGATCCT ATACTGCCAG AATACGAAGA AGAATACGTA GAAATTGAAA
GGATACAAGT TTTTACTTCT TACATGTCTA TGTCTAGGA TATGACGGTC TTATGCTTCT TCTTATGCAT CTTTAACTTT
401 AAGAAGAACC AGGCGAAGAA AAGAATCTTG AAGACGTAAG CACTGACGAC AACAAATGAAA AGAAGAAGAT AAGGTCGGTG
TTCTTCTTGG TCCGCTTCTT TTCTTAGAAC TTCIGCATTC GTGACTGCTG TTGTTACTTT TCTTCTTCTA TTCAGCCAC

FIG. 6A



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PstI

481 ATTGTGAAG AGACATAGAG GACACATGTA AGGTGGA AAA TGTAAGGGCT GCAGAAGGTA ATTATCCAAG ATGTAGCATC
TAACACTTTC TCTGTATCTC CTGTGTACAT TCCACCTTTT ACATCCCCGA CGTCTTCCAT TAATAGGTTT TACATCGTAG

SacI

561 AAGAAATCCAA TGTTACGGG AAAAATAATG GAAGTATTAT GTGAGCTCAG CAAGAAGCAG ATCAATAATGC GGCACATAATG
TTCTTAGGTT ACAAATGCC TTTTGATAC CTTTATAATA CACTCGAGTC GTTCTTCGTC TAGTTATACG CCGTGTATAC

641 CAACCTAATG TCAAAAATGA AGAATGTACA GATACAAGAT CCTATACTGC CAGAATACGA AGAAGAATAC GTAGAAATG
GTTGGATACA AGTTTTTACT TCTTACATGT CTATGTTCTA GGATATGACG GTCTTATGCT TCTTCTTATG CATCTTTAAC

721 AAAAAGAAGA ACCAGGCGAA GAAAAGAATC TTGAAGACGT AAGCACTGAC GACAACAATG AAAAGAAGAA GATAAGGTG
TTTTTCTTCT TGGTCCGCTT CTTTCTTAG AACTCTGCA TTCGTGACTG CTGTTGTTAC TTTTCTTCTT CTATCCAGC

801 GTGATTGTGA AAGAGACATA GAGGACACAT GTAAGGTGGA AAATGTAAGG GCGGAAAGTA ACCTTATCAC AAAGGAATCT
CACTAACACT TTCTCIGTAT CTCCTGTGTA CATTCCACCT TTACATTTCC CGCCTTTTAT TGGAAATAGTG TTTCTTTAGA

881 TATCCCCCAC TACTTATCCT TTTATATTTT TCCGTGTCAT TTTTGCCCTT GAGTTTCTCT ATATAAGGAA CCAAGTTCCG
ATAGGGGGTG ATGAATAGGA AAATATAAAA AGGCACAGTA AAAACGGGAA CTCAAAAGGA TATATTTCTT GGTTCAGGCC

961 CATTGTGTA AACAGA AAAA AATTGGTGT AAGCTATTTT CTTTGAAGTA CTGAGGATAC AACTTCAGAG AAATTGTGTA
GTAACACTT TTGTTCTTTT TTAACCCACA TTCGATAAAA GAACTTCAT GACTCCTATG TTGAAGTCTC TTTAACACTT

FIG. 6B

BamHI

1041 GTTTGTGGAT CC Seq. ID No. 5
 CAAACACCTA GG Seq. ID No. 6

FIG. 6C

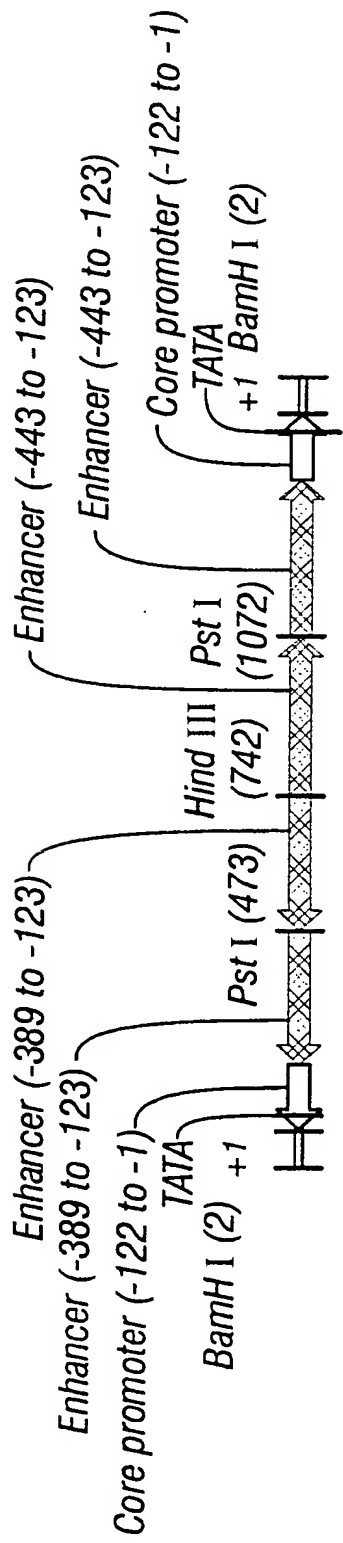


FIG. 7



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BamHI

1 GGATCCACAA ACTTACAAAT TTCTCTGAAG TTGTATCCTC AGTACTTCAA AGAAAATAGC TTACACCAAA TTTTCTCTTG
CCTAGGTGTT TGAATGTTTA AAGAGACTTC AACATAGGAG TCAIGAAGTT TCITTTATCG AATGIGGTTT AAAAAAGAAC

81 TTTTCACAAA TGCCGAACCT GGTTCCCTTAT ATAGGAAAAC TCAAGGGCAA AATGACACAG GAAAAATATA AAAGGATAAG
AAAAGTGTTC ACGGCTTGAA CCAAGGAATA TATCCTTTTG AGTTCCCGTT TTTACIGGC CTTTTTATAT TTTCTTATTC

161 TAGTGGGGGA TAAGATTCTT TTGTGATAAG GTTACTTTCC GCCCTTACAT TTCCACCTT ACAIGTGTCC TCTATGTCTC
ATCACCCCTT ATTCTAAGGA AACACTATTC CAATGAAAGG CGGGGATGTA AAAGTGGAA TGTACACAGG AGATACAGAG

241 TTTTCACAAT ACCGACCCTT TCTTCTCTT TTCAATTGTT TCGTCAGTGC TTACGCTTC AAGATTCCTT TCTTCGCCCTG
AAAGTGTAG TGGCTGGAAT AGAAGAAGAA AAGTAACAAC AGCAGTCAGG AATGCAGAAG TTCTAAGAAA AGAAGCGGAC

321 GTTCTTCTTT TTCAATTCT ACGTATCTT CTTCGTATTC TGGCAGTATA GGAICTTGT TCTGTACATT CTTCATTTT
CAAGAAGAAA AAGTTAAAGA TGCATAAGAA GAAGCATAAG ACCGTCATAT CCTAGAACAT AGACATGTAA GAAGTAAAAA

401 GAACATAGGT TGCATATGTG CCGCATATTG ATCTGCTTCT TGCTGAGCTC ACATAATACT TCCATAGCTG CAGCCCTTAC
CTTGATCCA ACGTATACAC GCGGTATAAC TAGACGAAGA ACGACTCGAG TGTATTAGA AGGATCGAC GTCGGGAATG

481 ATTTTCCACC TTACATGTGT CCTCTATGTC TCTTTCACAA TCACCGACCT TATCTTCTTC TTTTCATTGT TGTGTCAGT
TAAAAGGTGG AATGTACACA GGAGATACAG AGAAAGTGTT AGTGGCTGGA ATAGAAGAAG AAAAGTAACA ACAGCAGTCA

SacI PstI

FIG. 8A



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561 GCTIACGTCT TCAAGATTCT TTTCTTCGCC TGGTTCTTCT TTTTCAATTT CTACGTATTC TTCTTCGTAT TCIGGCAGTA
CGAATGCAGA AGTTCTAAGA AAGAAGCGG ACCAAGCGG AAGAGTAA GATGCATAAG AAGAAGCATA AGACCGTCAT

SacI

641 TAGGATCTTG TATCIGTACA TTCTTCATTT TTGAACATAG GTTGCATATG TGCCGCATAT TGATCTGCTT CTGTGAGC
ATCCTAGAAC ATAGACATGT AAGAAGTAAA AACTTGTAIC CAAGTATAC ACGCGTATA ACTAGACGAA GAACGACTCG

SacI

HindIII

721 TCACATAATA CTTCATAGG AAGCTTCAGA AGGTAATTAT CCAAGATGTA GCATCAAGAA TCCAATGTTT ACGGGAAGAA
AGTGTATTAT GAAGGATACC TTCGAAGICT TCCATTAAIA GGTTCTACAT CGTAGTTCTT AGGTIACAAA TGCCCTTTT

SacI

801 CTATGGAAGT ATTAIGTGAG CTCAGCAAGA AGCAGATCAA TATGCGGCAC ATATGCAACC TATGTTCAA AATGAAGAAT
GATACCTTCA TAATACACTC GAGTCGTCT TCGTCTAGTT ATACGCCGTG TATACGTTGG ATACAAGTTT TTACTTCTTA

881 GTACAGATAC AAGATCCTAT ACTGCCAGAA TACGAAGAAG AATACGTAGA AATTGAAAA GAAGAACCAG GCGAAGAAAA
CATGCTATG TTCTAGGATA TGACGGTCTT ATGCTTCTTC TTATGCATCT TTAACTTTTT CTTCCTGGTC CGCTTCTTT

961 GAATCTTGAA GACGTAAGCA CTGACGACAA CAATGAAAAA AAGAAGATAA GGTCGGTGAT TGTGAAAGAG ACATAGAGGA
CTTAGAACTT CTGCATTCTG GACTGCTGTT GTTACTTTTC TTCTTCTATT CCAGCCACTA ACACTTTCTC TGTATCTCCT

PstI

FIG. 8B



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1041 CACAIGTAAG GTGGAAAATG TAAGGGCTGC AGAAGGTAAT TATCCAAGAT GTAGCATCAA GAATCCAATG TTTACGGGAA
GTGTACATTC CACCTTTTAC ATTCCCGAGG TCTTCGTCTA ATAGGTCTA CATCGTAGT CTTAGGTIAC AAATGCCCTT

SacI

1121 AACTAATGGA AGTATTATGT GAGCTCAGCA AGAAGCAGAT CAATAIGCGG CACATAIGCA ACCTAIGTTC AAAAAATGAAG
TTTGATACCT TCATAATACA CTCGAGTCGT TCTTCGTCTA GTTATACGCC GTGTATACGT TGGATACAAG TTTTACTTC

1201 AATGTACAGA TACAAGATCC TATACTGCCA GAATACGAAG AAGAATACGT AGAAATTGAA AAAGAAGAAC CAGGCGAAGA
TTACATGCT ATGTTCTAGG ATATGACGGT CTTATGCTTC TTCTTATGCA TCTTAACTT TTTCTTCTTG GTCCGCTTCT

1281 AAAGAATCTT GAAGACGTAA GCACTGACGA CAACAATGAA AAGAAGAAGA TAAGGTGGT GATTGIGAAA GAGACATAGA
TTTCTTAGAA CTTCTGCATT CGTACTGCT GTTGTTACTT TTCTTCTTCT ATTCCAGCCA CTAACACTTT CTCTGTATCT

1361 GGACACATGT AAGGTGGAAA ATGTAAGGGC GGAAAGTAAC CTTATCACAA AGGAATCTTA TCCCCACTA CTTATCCTT
CCTGTGTACA TTCCACCCTT TACATTCCCG CCTTTCATTG GAATAGTGT TCCTTAGAAT AGGGGGTGAT GAATAGGAAA

1441 TATATTTTTC CGTGTCACTT TTGCCCTTGA GTTTTCTTAT ATAAGGAACC AAGTTCGGCA TTTGTGAAAA CAAGAAAAAA
ATATAAAAAG GCACAGTAAA AACGGGAAC CAAAGGATA TATTCCTTGG TTCAAGCCGT AACACCTTT GTTCTTTTT

BamHI

1521 TTITGGTGTA GCTATTTTCT TTGAAGTACT GAGGATACAA CTTCAGAGAA ATTTGTAAGT TTGTGGATCC Seq. ID No. 7
AAACCACATT CGATAAAAGA AACTTCATGA CTCCTATGTT GAAGTCTCTT TAAACATTCA AACACCTAGG Seq. ID No. 8

FIG. 8C

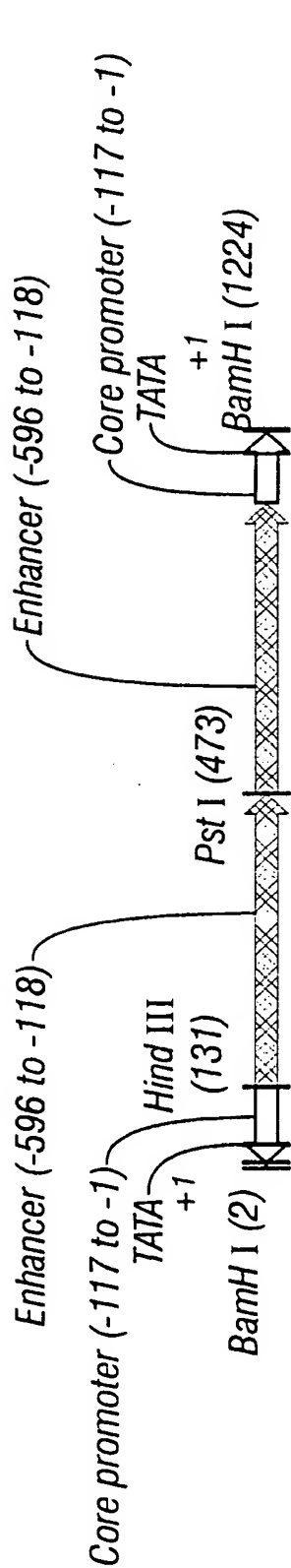


FIG. 9

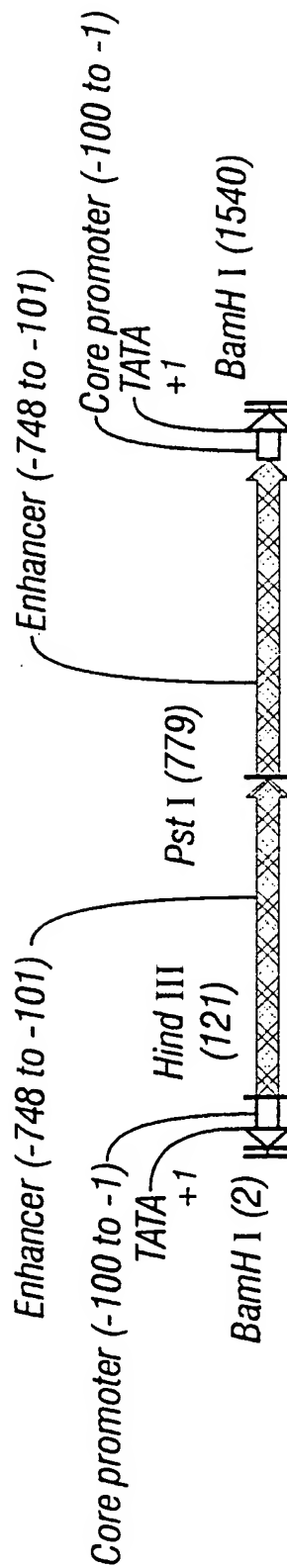


FIG. 11

BamHI

1 GGATCCTTGT TTTCAAAGCG GAGAGGAAAA TATATGAATT TATATAGCGG GGTTTATCTC TTACAACCTTT ATTTTCGGCC
CCTAGGAACA AAAGTTTCGC CTCCTCTTTT ATATACITAA ATATATCCGC CCAATAGAG AATGTTGAAA TAAAAGCCGG

HindIII

81 TTTCAAAAA ATAATTAAAA TCGACAGACA CGAATCATTT CGACCACAGA AGCTTCAACT ATTTTATGT ATGCAAGAGT
AAAGTTTTT TATTAATTT AGCTGCTGT GCTTAGTAAA GCTGGTGCT TCGAAGTTGA TAAAAATACA TACGTTCTCA

161 CAGCATATGT ATAATTGATT CAGAATCGTT TTGACGAGTT CGGATGTAGT AGTAGCCATT ATTAAATGTA CATACTAATC
GTCGTATACA TATTAACATA GTCTTAGCAA AACTGCTCAA GCCTACATCA TCATCGGTAA TAAATTACAT GTATGATTAG

241 GTGAATAGTG ATAATGATGAA ACATTGTATC TTATTGTATA AATATCCATA AACACATCAT GAAAGACACT TTCTTTCAG
CACATATCAC TATACTACTT TGTAAACATAG AATAACATAT TTATAGGTAT TTGIGTAGTA CTTTCTGTGA AAGAAAGTGC

321 GTCIGAATTA ATTAATGATAC AATTCTAATA GAAACGAAT TAAATTACGT TGAATTGTAT GAAATCTAAT TGAACAAGCC
CAGACTTAAT TAATACTATG TTAAGATTAT CTTTGTCTTA ATTTAATGCA ACTTAACATA CTTTAGATTA ACTTGTCGG

401 AACCACGACG ACGACTAACG TTGCCCTGGAT TGACTCGGTT TAAGTTAACC ACTAAAAAA CGGAGCTGTC ATGTAACACG
TTGGTGCTGC TGCTGATTGC AACGGACCTA ACTGAGCCAA ATTCAATTGG TGATTTTTT GCCTCGACAG TACATTGTGC

481 CGGATCGAGC AGGTCACAGT CATGAAGCCA TCAAGCAAA AGAACTAATC CAAGGCTGA GATGATTAAAT TAGTTTAAAA
GCCTAGCTCG TCCAGTGTCA GTACTTCGGT AGTTTCGTT TCTTGATTAG GTTCCCGACT CTACTAATTA ATCAAAATTT

PstI

FIG. 10A



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561 ATTAGTTAAC ACGAGGGAAA AGGCTGTCTG ACAGCCAGGT CACGTTAICT TTACCTGCAG CAACTAITTT TATGTATGCA
TAATCAATTG TGCTCCCTTT TCCGACAGAC TGTCGGTCCA GTGCAATAGA AATGGACGTC GTTGATAAAA ATACATACGT

641 AGAGTCAGCA TATGTATAAT TGATTCAGAA TCGTTTGGAC GAGTTCGGAT GTAGTAGTAG CCATTATTIA ATGTACATAC
TCTCAGTCGT ATACATATTA ACTAAGTCTT AGCAAAACTG CTCAAGCCTA CATCATCATC GGTAATAAAT TACATGTAIG

721 TAATCGTGAA TAGTGATATG ATGAAACATT GTATCTTATT GTATAAATAT CCATAAACAC ATCATGAAAG ACACTTTCTT
ATTAGCACTT ATCACTATAC TACTTTGTAA CATAGAATAA CATAITTIATA GGTATTIGTG TAGTACTTTC TGTGAAAGAA

801 TCACGGTCTG AATTAATTAT GATACAATTC TAATAGAAA CGAATTAAAT TACGTTGAAT TGTATGAAAT CTAATTGAAC
AGTGCCAGAC TTAATTAATA CTAIGTTAAG ATTAICTTTT GCITTAATTIA ATGCAACTTA ACATACITTA GATTAACITG

881 AAGCCAAACCA CGACGACGAC TAACGTTGCC TGGATTGACT CGGTTTAAGT TAACCACATAA AAAAAACGGAG CTGTCATGTA
TTCGGTIGGT GCTGCTGCTG ATTGCAACGG ACCTAACTGA GCCAAATTCA ATGGIGATT TTTTGGCCIC GACAGTACAT

961 ACACGCGGAT CGAGCAGGTC ACAGTCATGA AGCCATCAA GCAAAGAAC TAATCCAAGG GCTGAGATGA TTAATTAGTT
TGTGCGCCTA GCTCGTCCAG TGTCACTACT TCGGTAGTTT CGTTTCTIG ATTAGGTICC CGACTCTACT AATTAATCAA

1041 TAAAAATTAG TTAACACGAG GGAAAAGGCT GTCTGACAGC CAGGTCACGT TATCTTTACC TGTGGTCGAA ATGATTCGTG
ATTTTAAATC AATTGTGCTC CCTTTTCCGA CAGACTGTGG GTCCAGTGCA ATAGAAATGG ACACCAGCTT TACTAAGCAC

1121 TCTGTGCAAT TTAATTATTT TTTTGAAAGG CCGAAAATAA AGTTGTAAGA GATAAACCCG CCTATATAAA TTCATATATT
AGACAGCTAA AATTAATAAA AAAACTTCC GGCITTTATT TCAACATICT CTATTGGGC GGATATATTT AAGTATATAA

BamHI

FIG. 10B

1201 TTCCTCTCCG CTTTGAAAAC AAGGATCC Seq. ID No. 9
AAGGAGAGGC GAAACTTTTG TTCCTAGG Seq. ID No. 10

FIG. 10C

BamHI

1 GGATCCTTTT GGGTTTGGT GAGAAACAAG GAATAGTATG GATGGGTTTT AATAGGGAAT AAGAGTTGAA AAGTCTGCAA
CCTAGGAAA CCCAAAACCA CTCCTTGTTC CTTATCATAC CTACCCAAA TTAICCCCTTA TTCICAACCT TTCAGACGGT

HindIII

81 TTTGTAAAG AAAAAATTG GAAAGTCACA TGTTAGCAGA AGCTTCAGAC TCATTAACTT AAAAGAAGAT ATAGACTCAT
AAACATTTTC TTTTTTAAC CTTTCAGTGT ACAATCGTCT TCGAAGTCTG AGTAATTGAA TTTTCTTCTA TATCTGAGTA

161 TAACTTAAA GAAGATATAG ATTCCAACAC AAGTTCAAAA TTCATAAACG TCAATCTTGG CTAAATTTCT GAACATCAAT
ATTGAATTTT CTTCATATC TAAGGTTGTG TTCAAGTTTT AAGTATTGC AGTTAGAACC GATTAAAGA CTGTGAGTTA

241 GCATTCCCTT AAAATATAGA TAATAAGTTA GGATGTTGTC ACTTCTTAA AGCATATTCC GACTGAGTCT GGTAGAATCT
CGTAAGGAAA TTTTATATCT ATTATTCAAT CCTACAACAG TGAAGAATT TCGTATAAGG CTGACTCAGA CCACTTTAGA

321 CATAAACTTT AGGCCTTATC TCTTCAATTA GGCAATTACT TACCTCCGCT CTACTTTAAG AAAATTCAT GGAGTACACC
GTATTGAAA TCCGGAATAG AGAAGTTAAT CCGTTAATGA ATGGAGGCGA GATGAAATTC TTTTAAGTTA CCTCATGTGG

FIG. 12A

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401 ATTATTAAGT TCATATAAAA ATAAAATTAT ATTAATTCTG TCTCTTGTTG GTTCGGCTCTA TCTTTTCTG TTTTCCIGCT
    TAATAATTCA AGTATAATTT TATTTTAATA TAATTAAGAC AGAGAACAAAC CAAGCGAGAT AGAAAAAGAC AAAAGGACGA

481 TCAACCATAA CATATACAAG AACTACATTT TCCAAGCTAG ATATATCTAA CATGACTGAC TTTGTAAAT TCTTTTGCCA
    AGTGGTATT GTATAIGTTC TTGATGTAAA AGGTCGAIC TATATAGATT GTACTGACTG AAACATTTAA AGAAAAACGGT

561 AGTTAAAGAA AAAAAATGAT GTTATCCAAA TAATAAGAG AAAGAGCCCT AATGAAAAAA ATGATTTACT ATTAGAGTTG
    TCAATTCTT TTTTITACIA CAATAGGTTT ATTAATTCIC TTCTCGGA TTACTTTTT TACTAAATGA TAATCTCAAC

641 TTCAGCTAAT CACAICAATT ATGGTTTTCA TCAAGTATGA CTAATGGCGG CTCTTATCTC AGCTGATGTG ACATTGAAAT
    AAGTCGATTA GTGTAGTTAA TACCAAAAGT AGTTCATACT GATTACCGCC GAGAAATAGAG TGCACIACAC TGTAACITTA

721 TCTTTGACTT TAACACTAAT GTCATAATGCT TTCAAATTA TAATCCGATA AAGCTGCAGA CTCATTAACT TAAAAGAAGA
    AGAAACTGAA ATTGTGATTA CAGTATACGA AAGTTTAATT ATTAGGCTAT TTCGACGTCT GAGTAATTGA ATTTCTTCT

801 TATAGACTCA TTAACCTTAA AGAAGATATA GATTCCAACA CAAGTTCAA ATTCTATAAC GTCAATCTTG GCTAAATTC
    ATATCTGAGT AATTGAATTT TCTTCTATAT CTAAGGTGTG GTTCAAGTTT TAAGTATTG CAGTTAGAAC CGATTTAAAG

881 TGAACATCAA TGCATTCCCTT TAAATATAG ATAATAAGTT AGGATGTTGT CACTTCTTA AGCATATTC CGACTGAGTC
    ACTGTAGTT ACGTAAGGAA ATTTTATAIC TATTATTCAA TCCTACAACA GTGAAAGAAT TTCGTATAAG GCTGACTCAG

961 TGGTAGAATC TCATAAACTT TAGGCCTTAT CTCTTCAATT AGGCAATTAC TTACCTCCGC TCTACTTTAA GAAAATTCAA
    ACCATCTTAG AGTATTTGAA ATCCGGAATA GAGAAGTTAA TCCGTTAATG AATGGAGCGG AGATGAAAT CTTTTAAGTT
  
```

PstI

FIG. 12B

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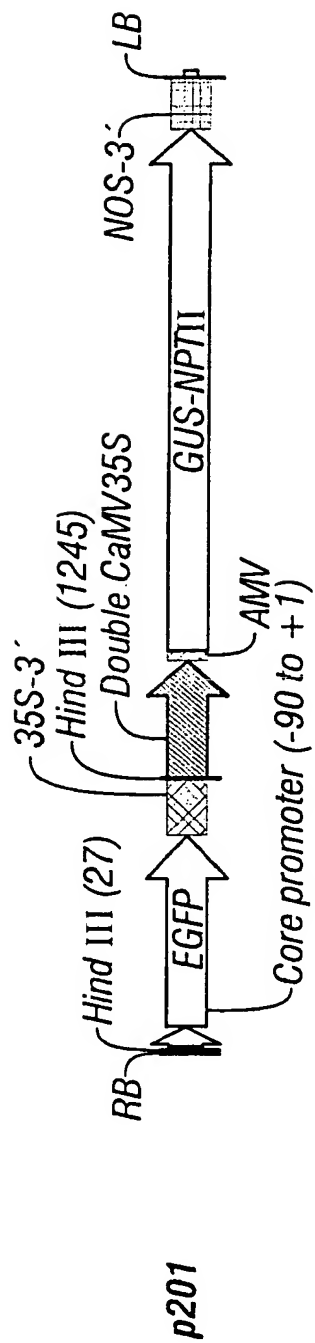


FIG. 13A

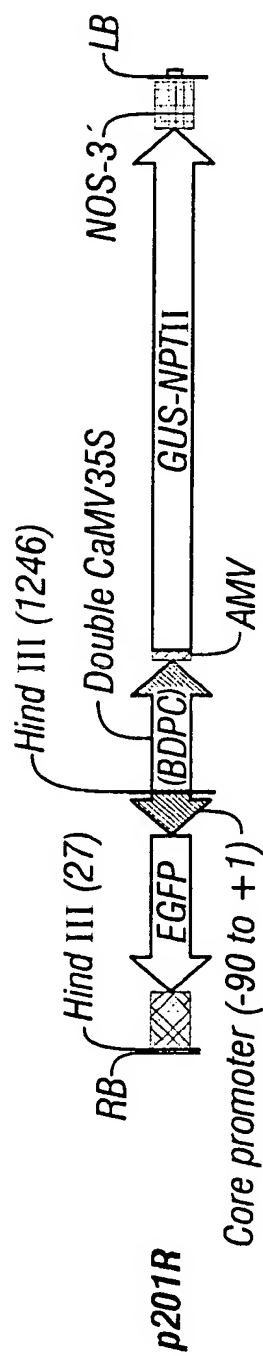


FIG. 13B



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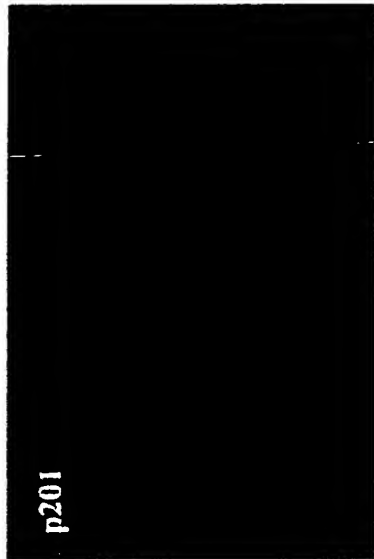


FIG. 14B

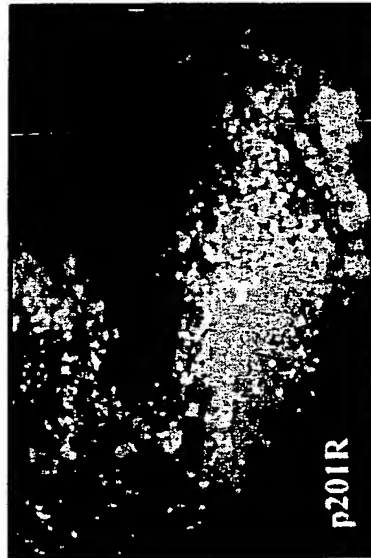


FIG. 14D

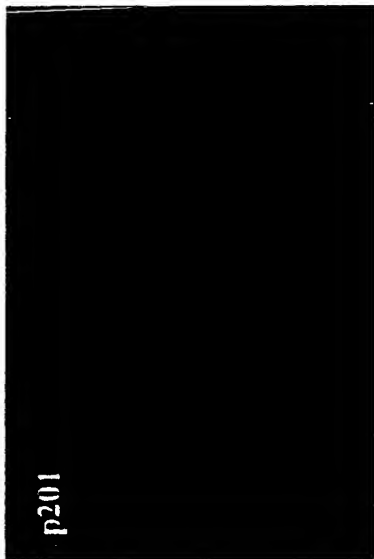


FIG. 14A

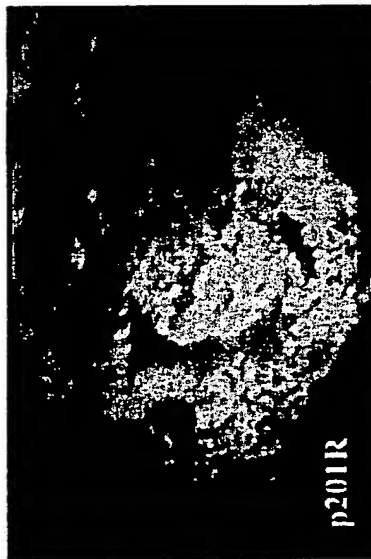


FIG. 14C

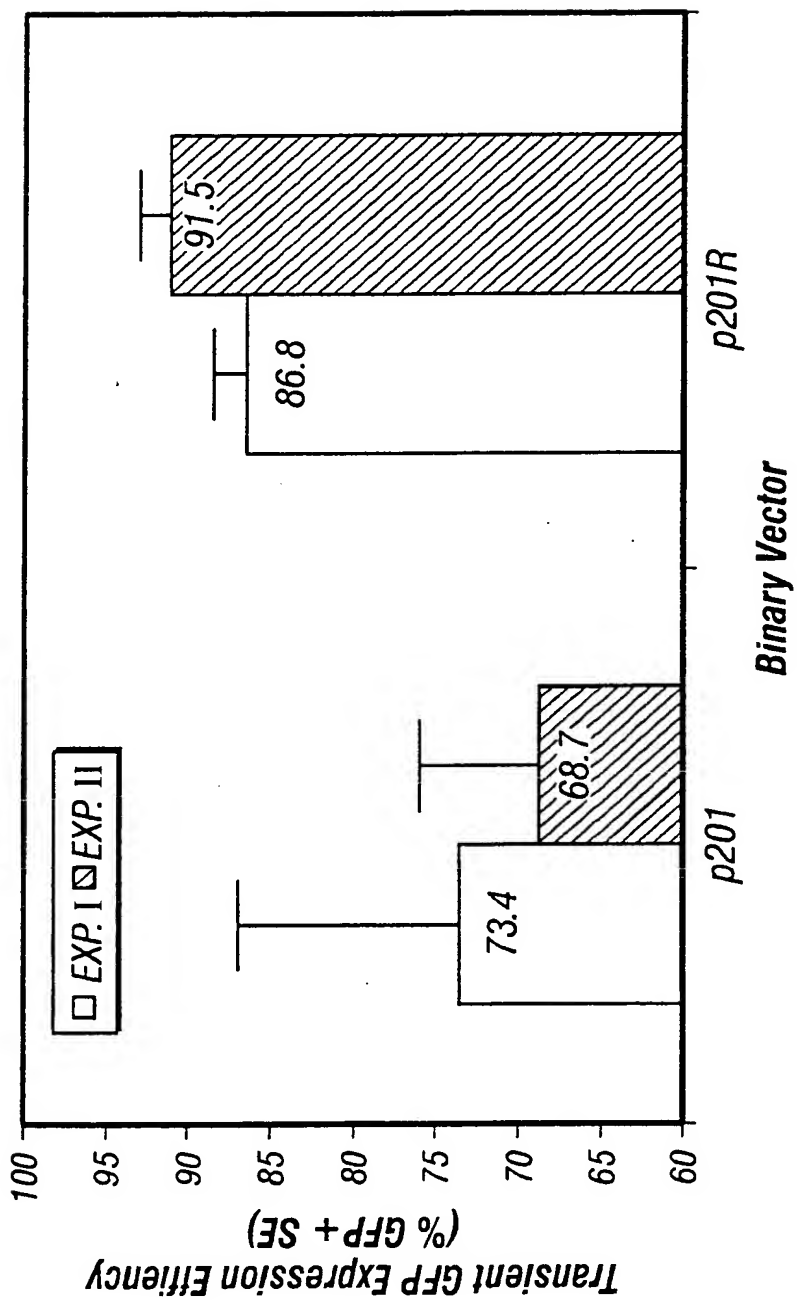


FIG. 15

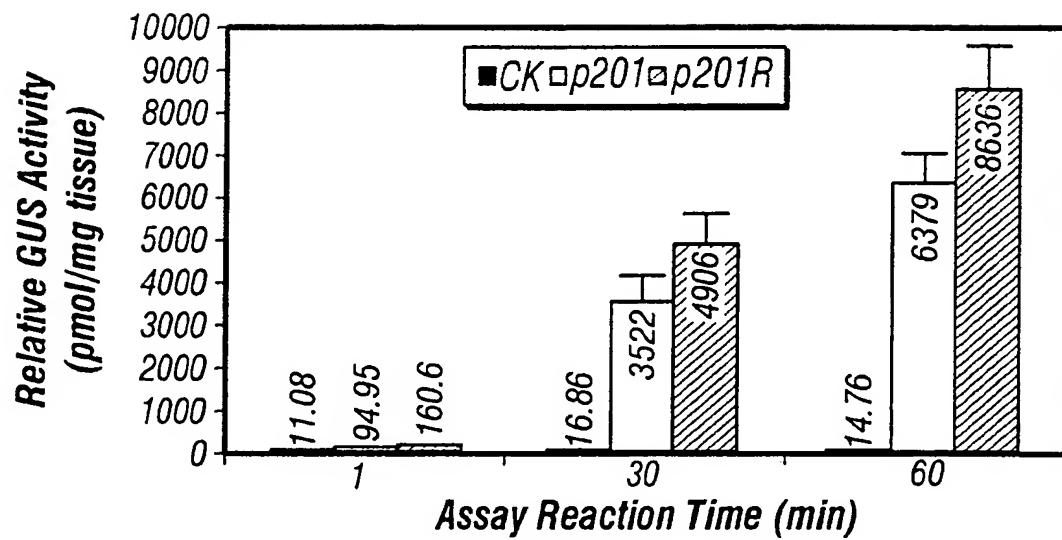


FIG. 16A

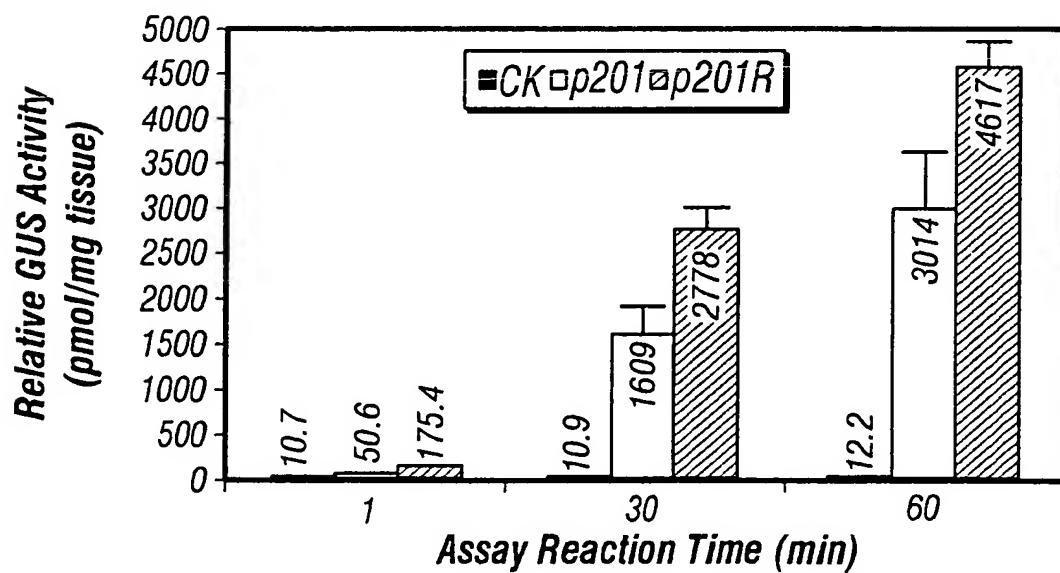


FIG. 16B

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FIG. 17A

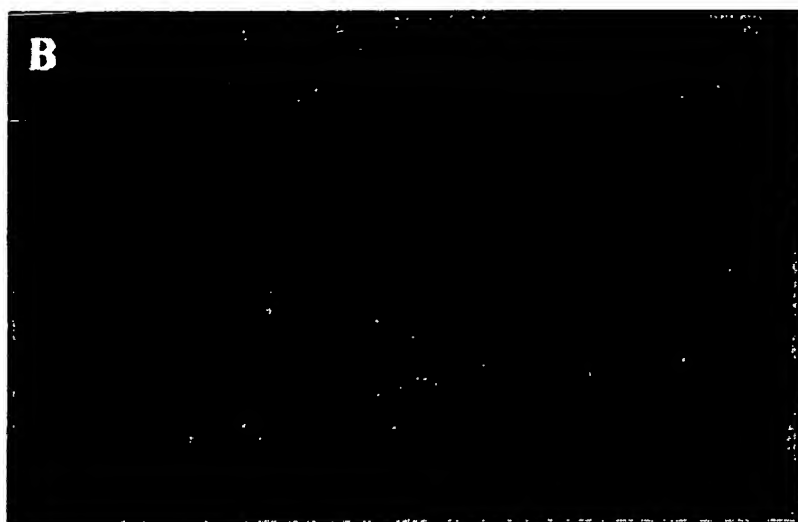


FIG. 17B

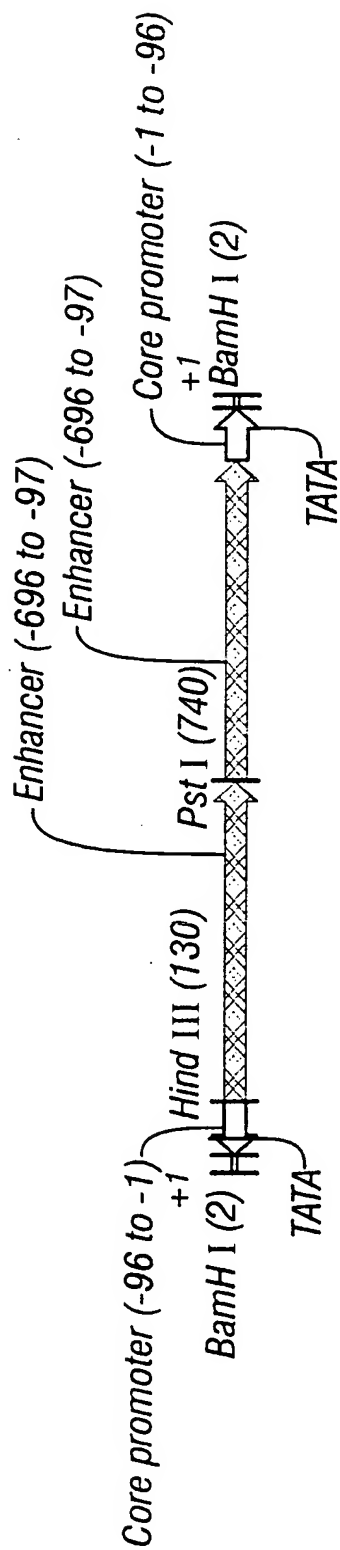


FIG. 18

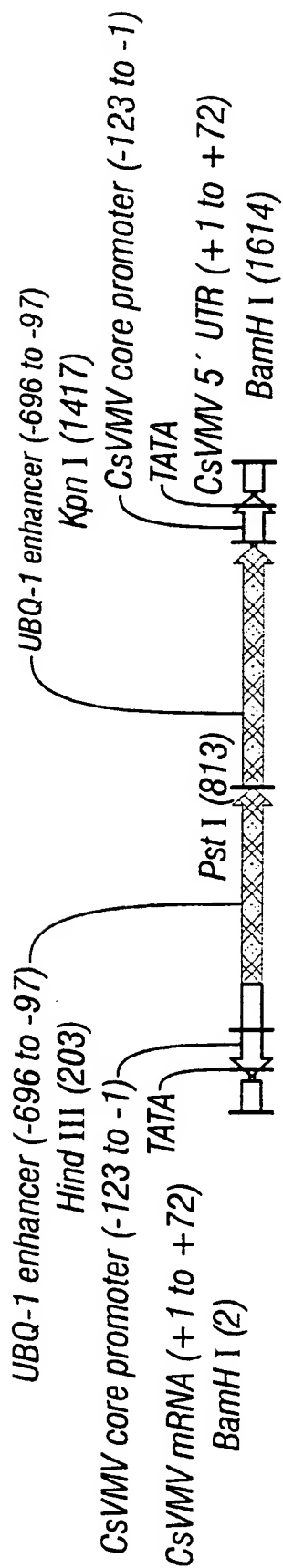


FIG. 20

BarHI

1 GGATCCCTTT TGTGTTTCGT CTTCTCTCAC GTAGAAACCC TAAACAAGGA GGAGGCGGGT TTATAATGT CAATGTACGC
CCTAGGGAAA ACACAAAGCA GAAGAGAGTG CATCTTTGGG ATTTGTTCTT CCTCCGCCCA AATATATACA GTTACATGCG

HindIII

81 GTCTAGGGTT TTGCTAATAT TGGGCTAGGT TACAGGCCTT TACCACAAAA GCTTAGTTGA TAAAAIATTT TTATTGGTT
CAGATCCCAA AACGATTATA ACCCGATCCA ATGTCCGGAA ATGGTGTTTT CGAATCAACT ATTTTATAA AATAAACCAA

161 GTAATTTTGT AATAICCCGG GATATTTCAC AAATTGAACA TAGACTACAG AATTTIAGAA ACAAACCTTT CTCICICTTA
CAITAAAACA TTATAGGGCC CTATAAGGCT TTTAACTTGT ATCTGATGTC TTAAATCTT TTGTTTGAAA GAGAGAGAAT

241 TCTCACCTTT ATCTTTTAGA GAGAAAAAGT TCGATTTCCG GTTGACCGGA ATGTAATCTT GTTTTTTTTG TTTTGTAAACA
AGAGTGGAAA TAGAAAAATCT CTCTTTTTCA AGCTAAAGC CAACTGGCCT TACATAGAAA CAAAAAAAC AAAACATTGT

321 TATTTCGTT TCCGATTIAG ATCGGATCTC CTTTCCGTT TTGTCGGACC TTCTCCGGT TTAICCGGAT CTAATAATAT
ATAAGCAAA AGGCTAAATC TAGCCTAGAG GAAAGGCCAA AACAGCCTGG AAGAAGGCCA AATAGGCCTA GATTATTATA

401 CCACTTTAGA CTTAGCTAAG TTGGATCTG TTTTTCGTT AGCTCTTGTC AATCGCCTCA TCAICAGCAA GAAGGTGAAA
GGTAGAATCT GAATCGATTG AACCTAGAC AAAAAACCAA TCGAGAACAG TTAGCGGAGT AGTAGTCGT CTTCACCTTT

481 TTTTGTACAA ATAAATCTTA GAATCATGTA GTGCTTTTGG ACCTTGGGAA TGATAGAAAC GATTGTGTAT AGCTACTCTA
AAAAACTGTT TATTTAGAAT CTTAGTACAT CACAGAAACC TGAACCCCTT ACTATCTTTG CTAACAATA TCGATGAGAT

FIG. 19A



561 TGTATCAGAC CCTGACCAAG ATCCAACAAT CTCATAGGTT TTGTGCATAT GAAACCTTCG ACTAACGAGA AGTGGTCITTT
ACATAGTCTG GGAAGTGGTC TAGGTGTTA GAGTATCCAA AACACGTATA CTTTGAAGC TGATTGCTCT TCACCAGAAA

641 TAATGAGAGA GATATCTAAA ATGTTATCTT AAAAGCCAC TCAAATCTCA AGGCATAAGG TAGAAATGCA AATTGGAAA
ATTACTCTCT CTATAGATTT TACAATAGAA TTTTCGGGTG AGTTTAGAGT TCCGTATTCC ATCTTTACGT TTAACCTTT

PstI
~~~~~

721 GTGGGCTGGG CCTTCTGCAG TTGATAAAAT ATTTTATTT GGTGTAAAT TTGTAATATC CCGGATATT TCACAAATTG  
CACCCGACCC GGAAGACGTC AACTATTTTA TAAAAATAA CCAACATTAA AACATTATAG GGCCCTATAA AGTGTTTAAC  
-----

801 AACATAGACT ACAGAAATTT AGAAACAAA CTTTCTCTCT CTTATCTCAC CTTTATCTTT TAGAGAGAAA AAGTTCGATT  
TTGTAICIGA TGTCTTAAAA TCTTTTGTG GAAAGAGAGA GAATAGAGTG GAAATAGAAA ATCTCTCTT TTCAAGCTAA  
-----

881 TCCGGTTGAC CGGAATGAT CTTTGTITTT TTGTTTGT AACATATTC GTTTCCGAT TTAGATCGGA TCTCCTTTTC  
AGGCCAACTG GCCTTACATA GAAACAAAAA AAACAAAAA TTGTATAAG CAAAAGGCTA AATCTAGCCT AGAGGAAAAAG  
-----

961 CGTTTGTGCG GACCTTCTTC CGGTTTATCC GGATCTAATA ATAICCATCT TAGACTTAGC TAAGTTTGGG TCTGTTTTT  
GCAAAACAGC CTGGAAGAAG GCCAAATAGG CCTAGATTAT TATAGGTAGA ATCTGAATCG ATTCAAACCT AGACAAAAA  
-----

1041 GGTTAGCCT TGTCAATCGC CTCATCATCA GCAAGAAGT GAAATTTTG ACAAATAAAT CTTAGAATCA TGTAGTCT  
CCAATCGAGA ACAGTTAGCG GAGTAGTAGT CGTCTTCCA CTTTAAAAAC TGTTTATTA GAATCTTAGT ACATCACAGA  
-----

1121 TTGGACCTTG GGAATGATAG AACGATTIG TTATAGCTAC TCTATGTATC AGACCTGAC CAAGATCCAA CAATCTCATA  
AACCTGGAAC CCTTACTATC TTTGCTAAAC AATATCGATG AGATACATAG TCTGGGACTG GTTCTAGGT GTTAGAGTAT  
-----

FIG. 19B



1201 GGTGTTGTGC ATAAGAAACC TTCGACTAAC GAGAAGTGGT CTTTAAATGA GAGAGATATC TAAATGTGA TCCTAAAAGC  
CCAAACACG TACTCTTTGG AAGCTGATTG CTCTTCACCA GAAATTAAT CTCTCTATAG ATTTTACAAT AGAATTTTCG

1281 CCACTCAAAT CTCAAGGCAT AAGGTAGAAA TGCAATTGG GAAAGTGGC TGGGCCTTTT GTGGTAAAGG CCIGTAACTT  
GGTGAGTTTA GAGTTCCGTA TTCCATCTTT ACGTTTAAAC CTTTCACCG ACCCGGAAAA CACCATTTCC GGACATTGGA

1361 AGCCCAATAT TAGCAAAACC CTAGACGCGT ACATTGACAT ATATAAACC GCCTCCTCCT TGTTTAGGGT TTCTACGTGA  
TCGGGTTATA ATCGTTTTGG GATCTGCGCA TGTAACCTGA TATATTTGG CGGAGGAGGA ACAAATCCCA AAGATGCACT

BamHI

1441 GAGAAGACGA AACACAAAAG GATCC Seq. ID No. 13

CTCTTCTGCT TTGTGTTTTTCT CTAGG Seq. ID No. 14

FIG. 19C

BamHI

1 GGATCCACAA ACTTACAAAT TTCCTGAAG TTGTATCCTC AGTACTTCAA AGAAATAGC TTACACCATA TTTTCTCTTG  
CCTAGGTGTT TGAATGTTTA AAGAGACTTC AACATAGGAG TCATGAAGTT TCCTTTATCG AATGTGGTTT AAAAAAGAAC

81 TTTTCACAAA TGCCGAACCT GTTTCCTTAT ATAGGAAAAC TCAAGGGCAA AAATGACACG GAAAAATATA AAAGGATAAG  
AAAAGTGTTC ACGGCTTGAA CCAAGGAATA TATCCTTTTG AGTTCCCGTT TTTACTGTGC CTTTCTATAT TTTCTTATTC

FIG. 21A



HindIII

161 TAGTGGGGA TAAGATTCCT TTGTGATAAG GTTACTTTCC GAAGCTTAGT TGATAAAATA TTTTATTG GTTGTAATTT  
ATCACCCCCT ATTCTAAGGA AACACTATTC CAATGAAAGG CTTGCAATCA ACTATTTTAT AAAAATAAAC CAACATTAAA  
-----  
241 TGTAAATACC CGGATATTT CACAAATTGA ACATAGACTA CAGAATTTTA GAAACAAAC TTTCTCTC TTATCTCACC  
ACATTATAGG GCCCTATAAA GIGTTAACT TGTATCTGAT GTCTTAAAG CTTTGTGTTG AAAGAGAGAG AATAGAGTGG  
-----  
321 TTTATCTTT AGAGAGAAA AGTTCGATT CCGGTGACC GGAATGTATC TTTGTTTTT TTGTTTGTG ACATATTTCG  
AAATAGAAA TCTCTCTTT TCAAGCTAAA GGCCAACCTGG CCTTACATAG AAACAAAAA AACAAAACAT TGTATAAGC  
-----  
401 TTTCCGATT TAGATCGGAT CTCCTTTTCC GTTTGTGCG ACCCTCTTCC GGTTTATCCG GATCTAATAA TATCCATCTT  
AAAAGGCTAA ATCTAGCCTA GAGGAAAAGG CAAACAGCC TGAAGAAGG CCAAATAGGC CTAGATTATT ATAGGTAGAA  
-----  
481 AGACTTAGCT AAGTTGGAT CTGTTTTTG GTTAGCTCTT GTCAATCGCC TCATCATCAG CAAGAAGGTG AAATTTTGA  
TCTGAATCGA TTCAAACCTA GACAAAAAAC CAATCGAGAA CAGTTAGCGG AGTAGTAGTC GTTCTTCCAC TTTAAAAACT  
-----  
561 CAAATAAATC TTAGAATCAT GTAGTGCTT TGGACCTTGG GAATGATAGA AACGATTGT TATAGCTACT CTATGTAICA  
GTTTATTAG AATCTTAGTA CATCACAGAA ACCTGGAACC CTTACTATCT TTGCTAAACA ATATCGAIGA GATACATAGT  
-----  
641 GACCCTGACC AAGATCCAAC AATCTCATAG GTTTGTGCA TAAGAAACCT TCGACTAACG AGAAGTGGTC TTTTAATGAG  
CTGGGACTGG TTCTAGGTTG TTAGAGTATC CAAACACGT ATACTTTGGA AGCTGATTGC TCTCACCAG AAAATTACTC  
-----  
721 AGAGATAICT AAAATGTTAT CTTAAAGCC CACTCAATC TCAAGGCATA AGGTAGAAAT GCAAATTTGG AAAGTGGGCT  
TCTCTATAGA TTTTACAATA GAATTTTCGG GTGAGTTTAG AGTCCGTAT TCCATCTTTA CGTTTAAACC TTTCAACCCGA  
-----



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PstI

801 GGGCCTTCTG CAGTTGATAA AATAATTTTA TTTGGTTGTA ATTTGTAAAT ATCCGGGAT ATTTCACAA TTGAACATAG  
CCCGGAAGAC GTCAACTATT TTATAAAAT AAACCAACAT TAAACATTA TAGGGCCCTA TAAAGTGT TTAACTGTATC  
-----  
881 ACTACAGAAT TTTAGAAAAC AACCTTCTC TCTCTTATCT CACCTTTATC TTTTAGAGAG AAAAGTTTCG ATTCCGGTT  
TGATGCTTA AAATCTTTTG TTTGAAAGAG AGAGAAATAGA GTGGAATAG AAAATCTCTC TTTTCAAGC TAAAGGCCAA  
-----  
961 GACCGGAATG TATCTTTGTT TTTTGTGTTT TGTAACATAT TTCGTTTTCC GATTAGATC GGATCTCCTT TTCCGTTTTC  
CTGGCCTTAC ATAGAAACAA AAAAAACAA ACATTGTATA AAGCAAAAGG CTAAATCTAG CCTAGAGGAA AAGGCAAAAC  
-----  
1041 TCGGACCTTC TTCCGGTTTA TCCGGATCTA ATAAATCCA TCTTAGACTT AGCTAAGTTT GGATCTGTTT TTTGGTTAGC  
AGCCTGGAAG AAGGCCAAAT AGGCCTAGAT TATTATAGGT AGAATCTGAA TCGATTCAAA CCTAGACAA AAACCAATCG  
-----  
1121 TCTTGTCAT CGCCTCATCA TCAGCAAGAA GGTGAAATTT TTGACAAATA AATCTTAGAA TCATGTAGTG TCTTTGGACC  
AGAACAGTTA GCGGAGTAGT AGTCGTCTT CCACITTAAT AACTGTTTAT TTAGAATCTT AGTACATCAC AGAAACCTGG  
-----  
1201 TTGGGAATGA TAGAAACGAT TTGTTATAGC TACTCTATGT ATCAGACCCT GACCAAGATC CAACAATCTC ATAGGTTTTC  
AACCTTACT ATCTTTGCTA AACAAATATCG ATGAGATACA TAGTCTGGA CTGGTTCTAG GTTGTAGAG TAICCAAAAC  
-----  
1281 TGCATATGAA ACCTTCGACT AACGAGAAGT GGTCITTTAA TGAGAGAGAT ATCTAAATG TTAICTTAA AGCCCACTCA  
ACGTATACTT TGGGAAGCTGA TTGCTCTTCA CCAGAAAAT ACTCTCTA TAGATTTTAC AATAGAATT TCGGGTGAGT  
-----

KpnI

FIG. 21C

1361 AATCTCAAGG CATAAGGTAG AAATGCAAAAT TTGAAAGTG GGCTGGGCCT TGGTACCCGG AAAGTAACCT TATCACAAAG  
 TTAGAGTTCC GTATTCCAATC TTACGTTTA AACCTTTCAC CCGACCCGGA ACCATGGGCC TTTCATTGGA ATAGTGTTTC

1441 GAATCTTATC CCCCACTACT TATCCTTTTA TATTTTCCG TGTCATTTT GCCCTTGAGT TTTCCTATAT AAGGAAGGAA  
 CTTAGAATAG GGGGTGATGA ATAGGAAAT ATAAAAGGC ACAGTAAAA CGGGAACCTCA AAAGGATATA TTCCTTGGTT

1521 GTTCGGCATT TGTGAAACA AGAAAAAAT TGGGTAGC TATTTCTTT GAAGTACTGA GGATACAACT TCAGAGAAAT  
 CAAGCCGTAA ACACCTTTGT TCTTTTAA ACCACATTGG ATAAAAGAAA CTTATGACT CCTATGTTGA AGTCTCTTTA

BamHI

1601 TTGTAAGTTT GTGGATCC Seq. ID No. 15  
 AACATTCAAA CACCTAGG Seq. ID No. 16

FIG. 21D

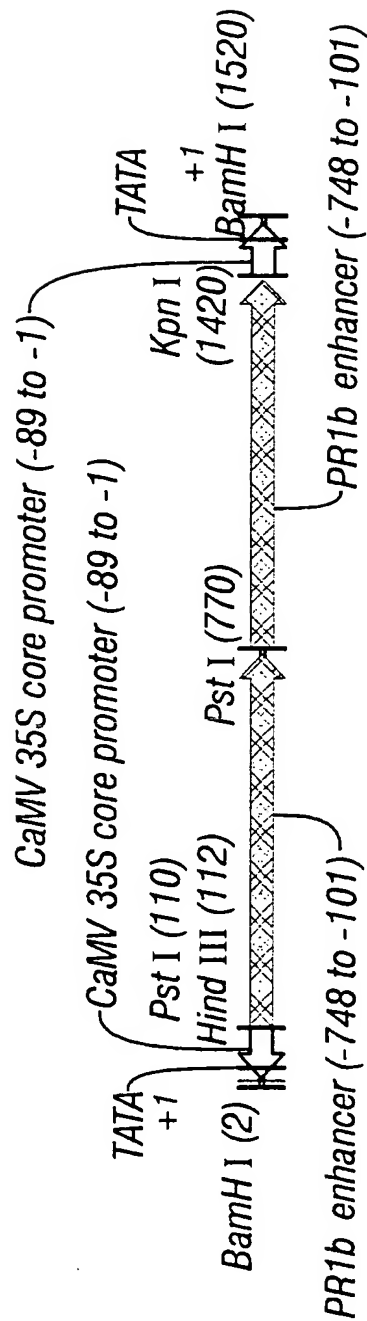


FIG. 22



BamHI

1 GGATCCAGCG TGTCCTCTCC AAATGAAATG AACTTCCTTA TATAGAGGAA GGGTCTTGGC AAGGATAGTG GGATTGTGCG  
CCTAGTGC ACAGGAGAGG TTTACTTTAC TTGAAGGAAT ATATCTCCTT CCCAGAAGCG TTCCTATCAC CCTAACACGC

PstI HindIII

81 TCATCCCTTA CGTCAGTGA GATACTGCAG AAGCTTCAGA CTCATTAACT TAAAGAAGA TATAGACTCA TTAACCTTAA  
AGTAGGGAAT GCAGTCACCT CTATGACGTC TTCGAAGTCT GAGTAATTGA ATTTCTTCT ATATCTGAGT AATTGAATTT

161 AGAAGATATA GATTCCAACA CAAGTTCANA ATTCATAAAC GTCAATCTTG GCTAAATTC TGAACATCAA TGCATTCCTT  
TCTTCTAT CTAGGTTGT GTTCAAGTTT TAAGTATTG CAGTTAGAAC CGATTTAAG ACTTGAGT ACGTAAGGAA

241 TAAATATAG ATAATAAGTT AGGATGTTGT CACTTCTTA AAGCATATTC CGACTGAGTC TGGTAGAATC TCATAAAGTT  
ATTTATATC TATTATTCAA TCCTACAACA GTGAAGAAT TTCGTATAAG GCTGACTCAG ACCATCTTAG AGTATTGAA

321 TAGGCCTTAT CTCTTCAATT AGGCAATTAC TTACCTCCGC TCTACTTTAA GAAAATTCAC TGGAGTACAC CATTATTAAAG  
ATCCGGAATA GAGAAGTTAA TCCGTTAATG AATGGAGGCG AGAIGAAAT CTTTAAAGT ACCTCATGIG GTAATAATTC

401 TTCATATAAA AATAAAATTA TATTAATTCT GTCTCTTGT GGTTCGCTCT ATCTTTTCT GTTTTCTGCTTCAACCATTA  
AAGTATAATT TTATTTAAT ATAATTAAGA CAGAGAACAA CCAAGCGAGA TAGAAAAAGA CAAAAGGAGC AAGTTGGTAT

481 ACATATACAA GAACTACATT TTCCAAGCTA GATATATCTA ACATGACTGA CTTTGTAAT TTCTTTTGCC AAGTTAAAGA  
TGATAIGTT CTTGATGTA AAGGTTGAT CTATATAGAT TGTACTGACT GAAACATTAA AAAAAAGCC TTCAATTTCT

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FIG. 23A





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561 AAAAAATGA TGTATCCAA ATAATAAGA GAAAGAGCCC TAATGAAAA AATGATTAC TATTAGATT GTTCAGCTAA  
TTTTTTACT ACAATAGGT TATTATTCT CTTCTCGGG ATTACTTTT TTAATAAATG ATAATCTCAA CAAGTCGATT

641 TCACATCAAT TATGGTTTT ATCAAGTATG ACTAATGGCG GCCTTAICT CACGIGATGT GACATTGAAA TTCTTTGACT  
AGGTAGTTA ATACCAAAG TAGTTCATAC TGATTACCGC CGAGAATAGA GTGCACTACA CTGTAACCTT AAGAAACTGA

PstI

721 TTAACACTAA TGTCATATGC TTTCAAATTA ATAATCCGAT AAAGCTGCAG ACTCATTAAC TTAAAAGAAG ATATAGACTC  
AATTGIGATT ACAGTATACG AAAGTTTAAT TATTAGGCTA TTTCGACGTC TGAGTAATTG AATTTTCTTC TATATCTGAG

801 ATTAACCTAA AAGAAGATAT AGATTCCAAC ACAAGTTCAA AATTCATAAA CGTCAATCTT GGCTAAATTT CTGAACATCA  
TAATTGAATT TTCTTCTATA TCTAAGGTG TGTTCAGTT TTAAGTATT GCAGTAGAA CCGATTTAAA GACTTGTAGT

881 ATGCATTCCT TTAAATATA GATAATAAGT TAGGATGTTG TCACTTTCTT AAAGCATATT CCGACTGAGT CTGGTAGAAT  
TAGCTAAGGA AATTTTATAT CTATTATCA ATCCTACAAC AGTGAAGAA TTTCGTATAA GGCTGACTCA GACCATCTTA

961 CTCATAAACT TTAGGCCCTTA TCTCTTCAAT TAGGCAATTA CTTACCTCCG CTCTACTTTA AGAAAAATCA ATGGAGTACA  
GAGTATTGA AATCCGGAAT AGAGAAGTTA ATCCGTTAAT GAATGGAGGC GAGATGAAAT TCTTTTAAGT TACCICATGT

1041 CCATTATTAA GTTCATAATA AAATAAAAT ATATTAATC TGTCCTGCTC TATCTTTTTC TGTTTTCCTG  
GGTAATAAT CAAGTATATT TTTATTTTAA TATAATTAAG ACAGAGAACA ACCAAGCGAG ATAGAAAAAG ACAAAGGAC

FIG. 23B



|      |                                                                                           |       |
|------|-------------------------------------------------------------------------------------------|-------|
| 1121 | CTTCAACCAT AACATATACA AGAACTACAT TTTCCAAGCT AGATAATCT AACATGACTG ACTTIGTAAA TTCTTTTTCG    |       |
|      | GAAGTTGGTA TTGTATACT TCTTGATGTA AAAGGTTTGA TCTATATAGA TTGTACTGAC TGAACACATTT AAAGAAAAACG  |       |
| 1201 | CAAGTTAAAG AAAAAAATG ATGTTATCCA AATAATAAG AGAAAGAGCC CTAATGAAAA AAATGATTTA CTATTAGAGT     |       |
|      | GTTCAATTTC TTTTITTTAC TACAATAGGT TTATTATTTC TCTTCTCGG GATTACTTTT TTTACTAAAT GATAATCTCA    |       |
| 1281 | TGTTCAAGCTA ATCACAATCAA TTATGGTTTT CATCAAGTAT GACTAATGGC GGCCTTTATC TCAGGTGATG TGACATTGAA |       |
|      | ACAAGTCGAT TAGTGATGTT AATACCAAAA GTAGTTCATA CTGATTACCG CCGAGAATAG AGTGCACTAC ACTGTAACTT   |       |
| 1361 | ATTCTTTGAC TTTAACACTA ATGTCATATG CTTTCAAATT AATAATCCGA TAAAGGTACC TATCTCCACT GACGTAAGGG   |       |
|      | TAAGAACTG AAATTGIGAT TACAGTATAC GAAAGTTTAA TTATTAGGT ATTICCAITGG ATAGAGGTGA CTGCATTCCC    |       |
| 1441 | ATGACGCACA ATCCCACTAT CCTTCGCAAG ACCCTTCCTC TATATAAGGA AGTTCATTTC ATTTGGAGAG GACACGCTGG   |       |
|      | TACTGCGTGT TAGGTGATA GGAAGCGTTC TGGGAAGGAG ATATATTCTT TCAAGTAAAG TAAACCTCTC CTGTGCGACC    |       |
|      | BamH                                                                                      | BamHI |
| 1521 | ATCC                                                                                      |       |
|      | TAGG                                                                                      |       |

FIG. 23C

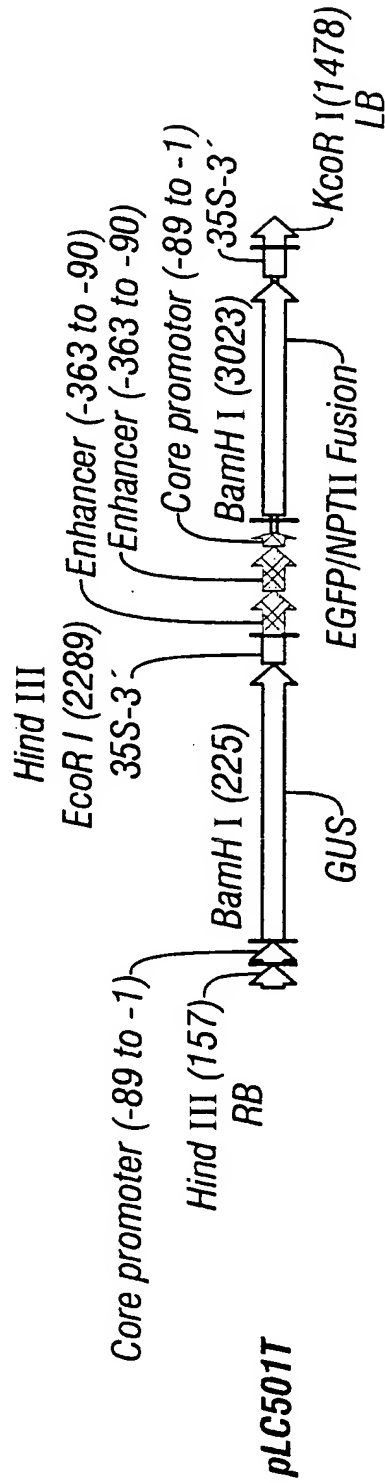


FIG. 24A

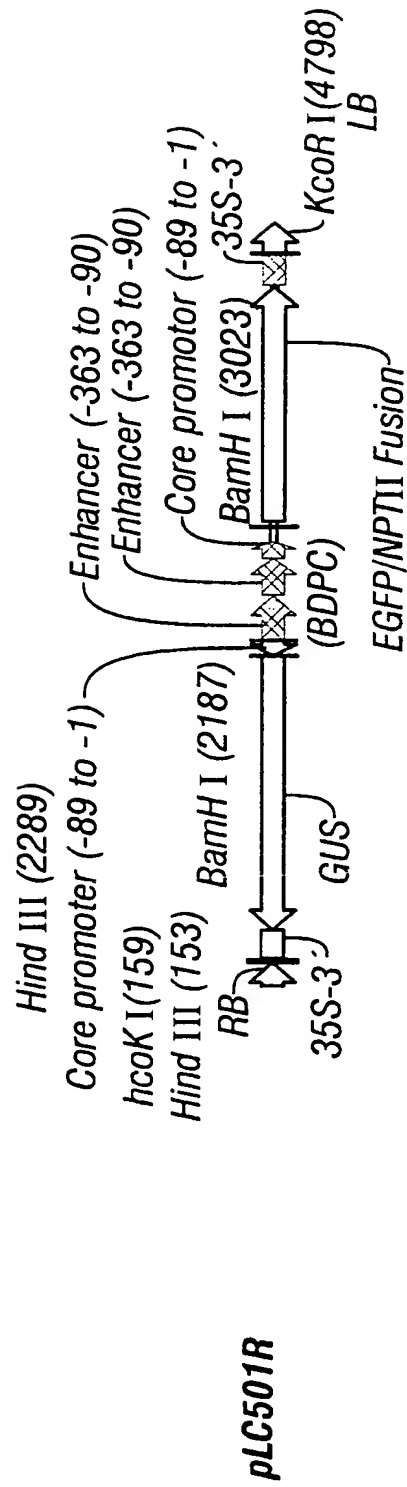


FIG. 24B

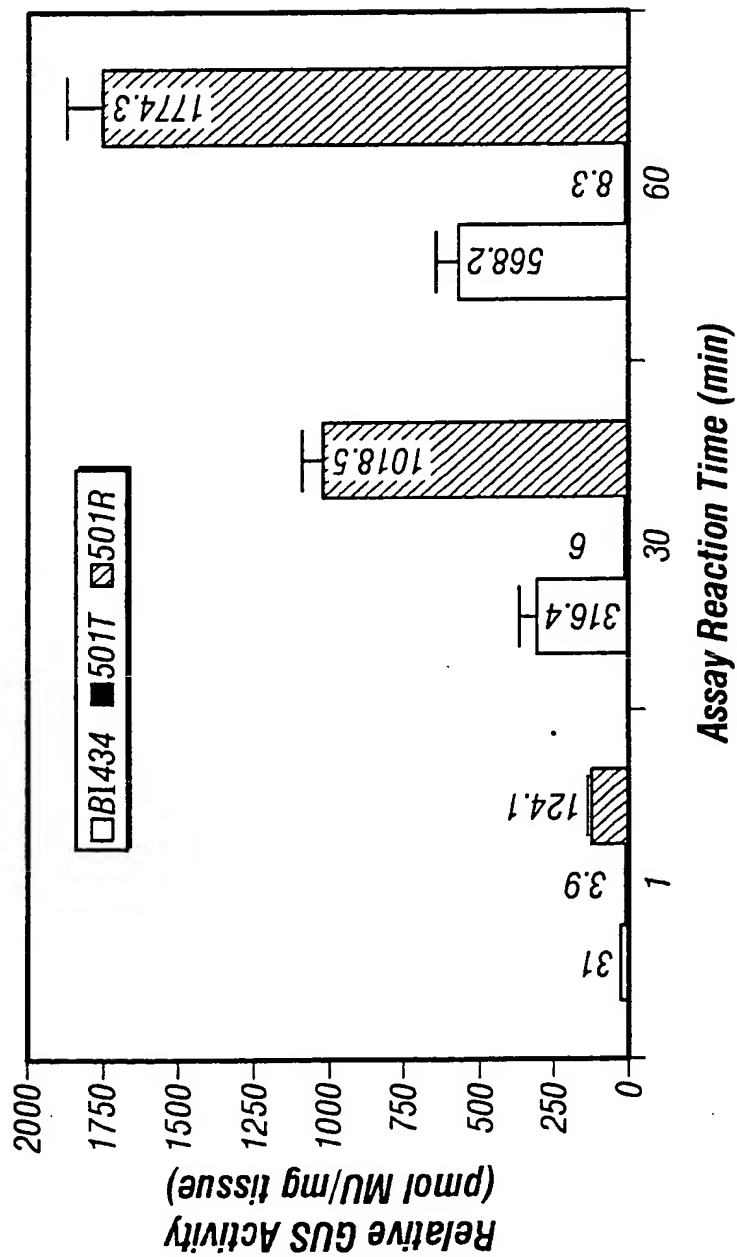


FIG. 25

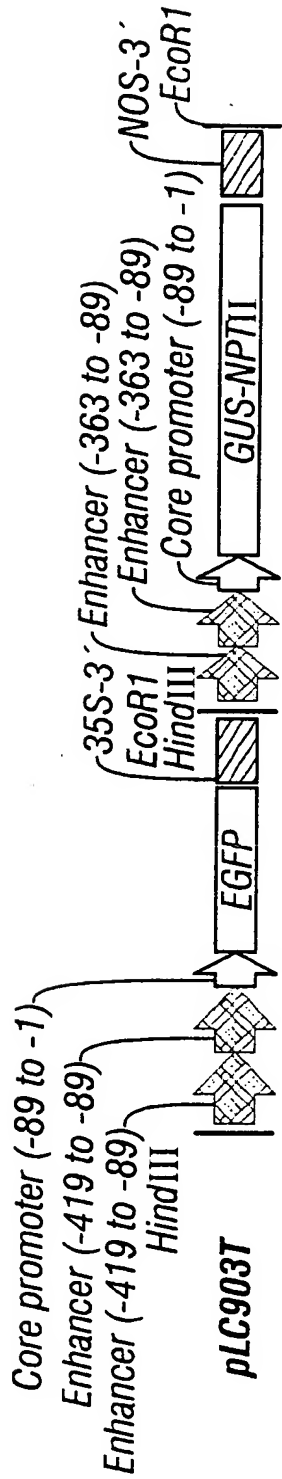


FIG. 26A

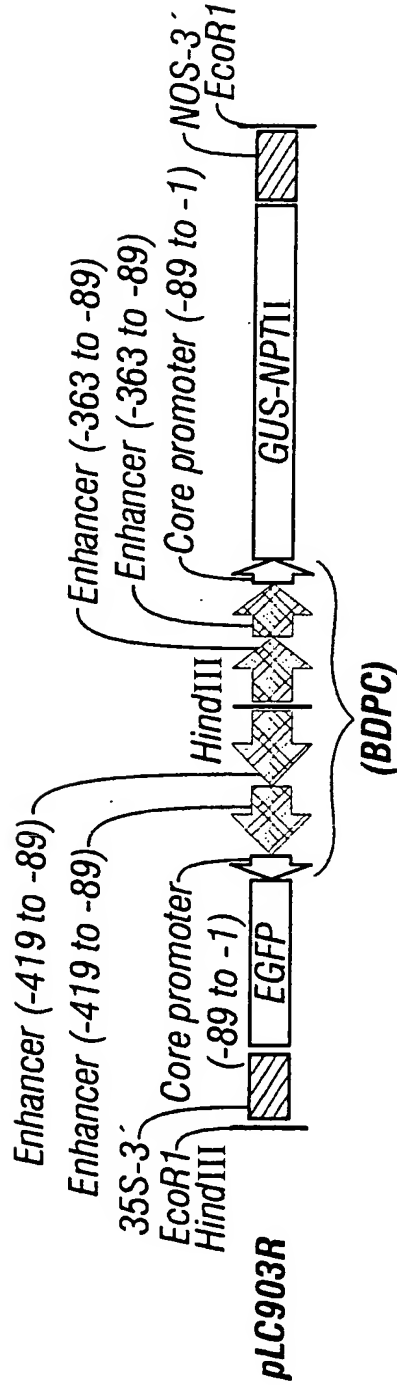


FIG. 26B



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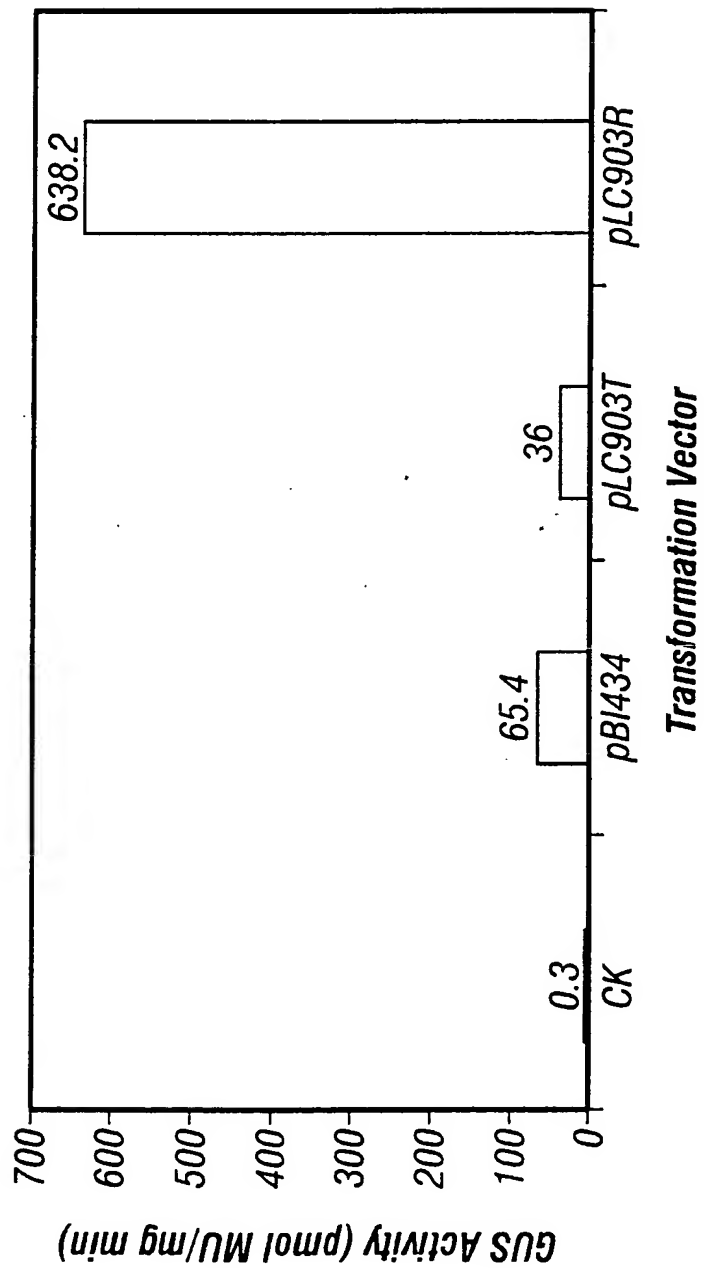


FIG. 27